

PF-0572 US

60 / 150,689

not hairy

# seq 2, 4

SEQUENCE LISTING

<110> Tang, Y. Tom  
Corley, Neil C.  
Patterson, Chandra

<120> HUMAN TAFAZZIN PROTEIN

<130> PF-0572 US

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<170> FastSEQ for Windows Version 3.0

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<212> PRT

<213> Homo Sapiens

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Glu	Pro	Gly	Asp	Gly	Asp	Arg	Glu	Met	Ala	Ser	Gly	Val	Gly	Gly	Leu
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Gly	Leu	Pro	Leu	Val	Pro	Gly	Cys	Pro	Ala	Pro	Pro	His	Val	Trp	Pro
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Ser	Val	His	Cys	Ala	Ala	Gly	Met	Asn	Asp	Val	Leu	Pro	Asn	Ser	Pro
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Pro	Tyr	Phe	Pro	Arg	Phe	Gly	Gln	Lys	Ile	Thr	Val	Leu	Ile	Gly	Lys
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Pro	Phe	Ser	Ala	Leu	Pro	Val	Leu	Glu	Arg	Leu	Arg	Ala	Glu	Asn	Lys
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Ser	Ala	Val	Glu	Met	Arg	Lys	Ala	Leu	Thr	Asp	Phe	Ile	Gln	Glu	Glu
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Phe	Gln	His	Leu	Lys	Thr	Gln	Ala	Glu	Gln	Leu	His	Asn	His	Leu	Gln
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Pro	Gly	Arg													
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35 40 45
Val Leu Tyr Glu Leu Ile Glu Lys Arg Gly Pro Ala Thr Pro Leu Ile
50 55 60
Thr Val Ser Asn His Gln Ser Cys Met Asp Asp Pro His Leu Trp Gly
65 70 75 80
Ile Leu Lys Leu Arg His Ile Trp Asn Leu Lys Leu Met Arg Trp Thr
85 90 95
Pro Ala Ala Ala Asp Ile Cys Phe Thr Lys Glu Leu His Ser His Phe
100 105 110
Phe Ser Leu Gly Lys Cys Val Pro Val Cys Arg Gly Ala Glu Phe Phe
115 120 125
Gln Ala Glu Asn Glu Gly Lys Gly Val Leu Asp Thr Gly Arg His Met
130 135 140
Pro Gly Ala Gly Lys Arg Arg Glu Lys Gly Asp Gly Val Tyr Gln Lys
145 150 155 160
Gly Met Asp Phe Ile Leu Glu Lys Leu Asn His Gly Asp Trp Val His
165 170 175
Ile Phe Pro Glu Gly Lys Val Asn Met Ser Ser Glu Phe Leu Arg Phe
180 185 190
Lys Trp Gly Ile Gly Arg Leu Ile Ala Glu Cys His Leu Asn Pro Ile
195 200 205
Ile Leu Pro Leu Trp His Val Gly Met Asn Asp Val Leu Pro Asn Ser

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210	215	220
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Lys Pro Phe Ser Ala Leu Pro Val Leu Glu Arg Leu Arg Ala Glu Asn		240
	245	250
Lys Ser Ala Val Glu Met Arg Lys Ala Leu Thr Asp Phe Ile Gln Glu		255
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Glu Phe Gln His Leu Lys Thr Gln Ala Glu Gln Leu His Asn His Leu		270
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Gln Pro Gly Arg		285
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35 40 45
Glu Ile Gln Lys Lys Leu Glu Ala Ala Glu Glu Arg Arg Lys Ser His
50 55 60
Glu Ala Glu Val Leu Lys Gln Leu Ala Glu Lys Arg Glu His Glu Lys
65 70 75 80
Glu Val Leu Gln Lys Ala Ile Glu Glu Asn Asn Asn Phe Ser Lys Met
85 90 95
Ala Glu Glu Lys Leu Thr His Lys Met Glu Ala Asn Lys Glu Asn Arg
100 105 110
Glu Ala Gln Met Ala Ala Lys Leu Glu Arg Leu Arg Glu Lys Asp Lys
115 120 125
His Ile Glu Glu Val Arg Lys Asn Lys Glu Ser Lys Asp Pro Ala Asp
130 135 140
Glu Thr Glu Ala Asp
145